

## Sep 7<sup>th</sup>

### -PCR Amplification of SmKSL1

[Volume] (50  $\mu$ L):

PrimerStar: 25  $\mu$ L

KSL1-Forward: 1.5  $\mu$ L

KSL1-Reverse: 1.5  $\mu$ L

Template (KSL1-32b, 10X): 1

$\mu$ L ddH<sub>2</sub>O: 21  $\mu$ L

[Program] (3 steps 34cycles):

98°C 2'

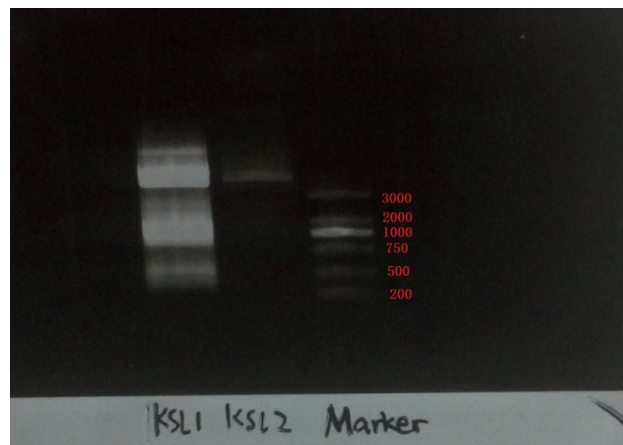
98°C 10''

60°C 5''

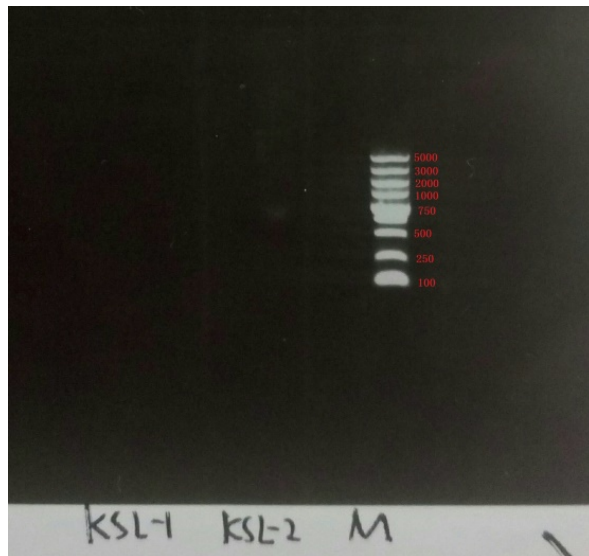
72°C 1'40''

72°C 5'

### -Results:



(failed)



(failed)

## Sep 8<sup>th</sup>

-PCR Amplification of SmKSL1

Program changed to:

98°C 2'

98°C 10''

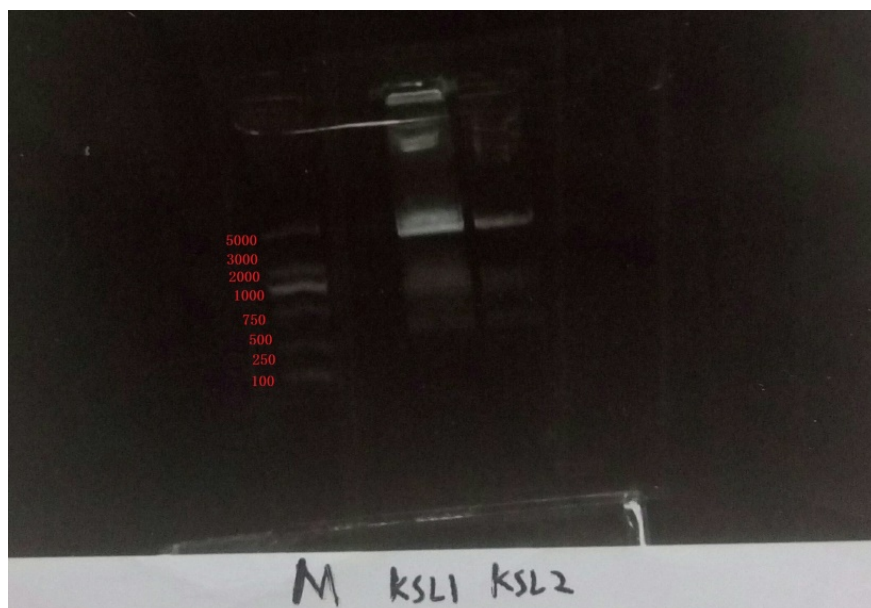
64°C 5''

72°C 2'

72°C 5'

34 cycles

-Result:



## Sep 9<sup>th</sup>

- Connect SmKSL1 to t-vector
- Transformation and spread plate method

## Sep 10<sup>th</sup>

- Two single colonies are selected for TA Cloning Test

## Sep 11<sup>th</sup>

- TA Cloning Test (Blunt-ended t-vector: pEASY- BluntZero Cloning Vector

[Volume](20  $\mu$ L):

Mix 10  $\mu$ L

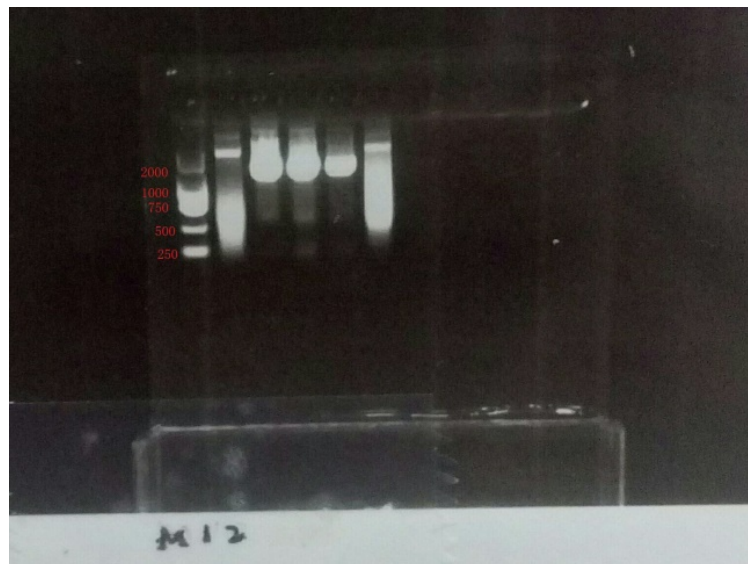
F 0.5  $\mu$ L

R 0.5  $\mu$ L

Template 0.5  $\mu$ L

ddH<sub>2</sub>O 8.5  $\mu$ L

- Result: (1-failed, 2-successful)



- Mutagenesis of Xba I (1225)

PCR [Volume] (50  $\mu$ L):

5xPrimeSTAR GXL Buffer 10  $\mu$ L

dNTP Mix (2.5mm each) 4  $\mu$ L

Primer1 1.5  $\mu$ L

Primer2 1.5  $\mu$ L

Template 0.5  $\mu$ L

PrimeSTAR GXL DNA Polymerase 1  $\mu$ L

ddH<sub>2</sub>O 31.5  $\mu$ L

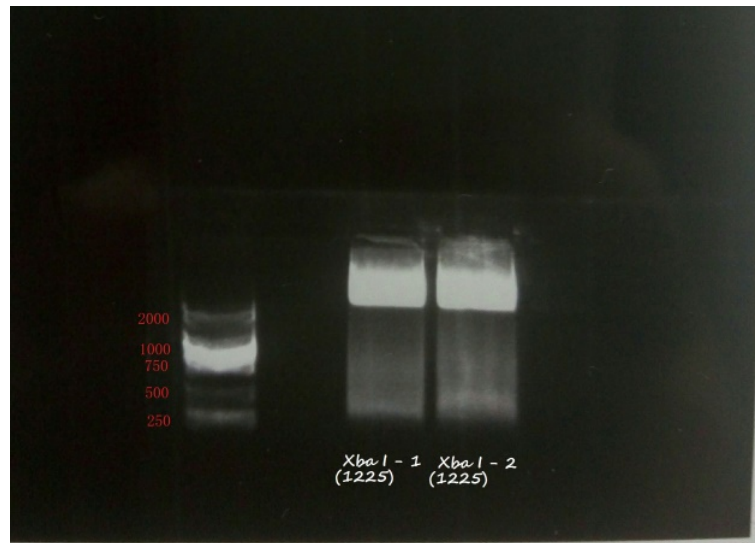
[Program] (30cycles, 3-step):

98°C 10''

62°C 15''

68°C 6min (1min/kb)

-PCR Result:



- Enzymatic digestion(enzyme: Takara 1235S Dpn I)

[Volume](20  $\mu$ L):

Dpn I 1  $\mu$ L

10x T Buffer 2  $\mu$ L

DNA 17  $\mu$ L

## Sep 12<sup>th</sup>

-Transformation & Spread Plate Method

## Sep 13<sup>th</sup>

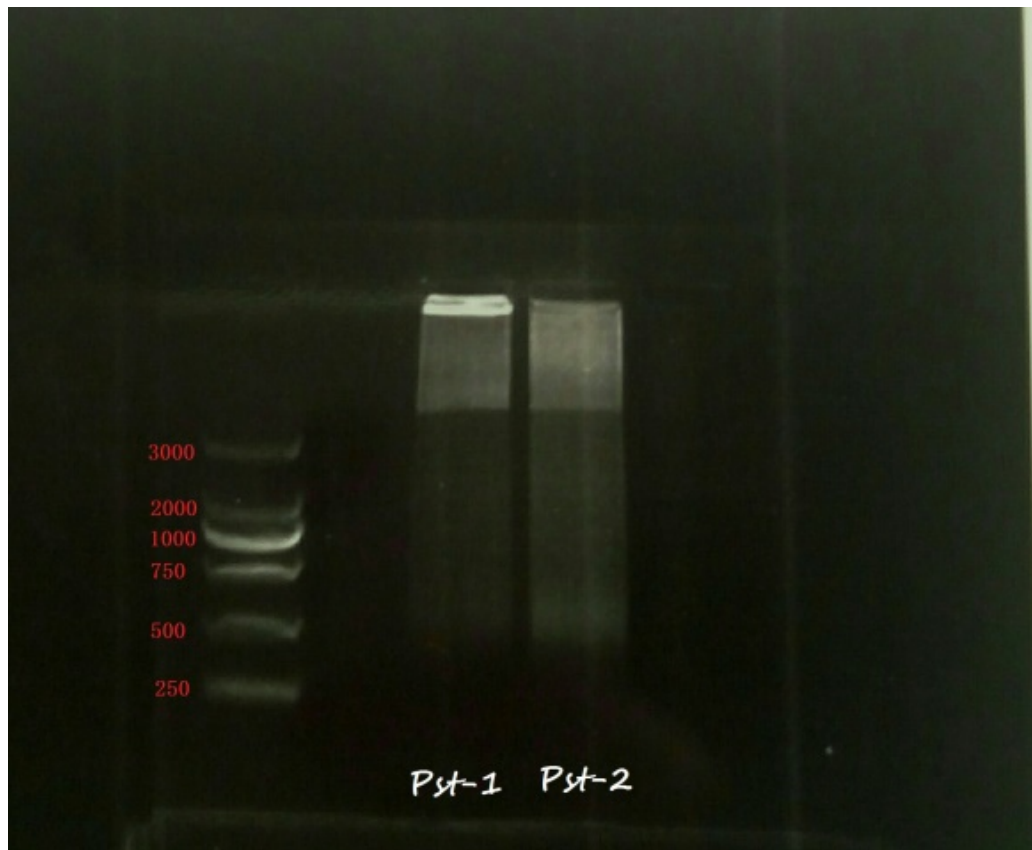
-Ten single colonies are selected for gene sequencing to check if the mutagenesis is managed

## Sep 14<sup>th</sup>

-Sequencing Result: No.4 & 7 successful

-Mutagenesis of Pst (1316)

PCR Result(unsuccesful)



**Sep 15<sup>th</sup>**

-Mutagenesis PCR of EcoR I (1688), Pst I (1316) and Xba I (322)

PCR Result: (Xba I (322) successful)



-Digestion, transformation, and spread plate method

### Sep 16<sup>th</sup>

- Ten single colonies are selected for gene sequencing to check if the mutagenesis is managed

### Sep 17<sup>th</sup>

-Sequencing results show that all colonies failed